



```

chain nodes :
 7 8 9 10 11 12 13 17 24 25 26 27 28 29 30 34
ring nodes :
 1 2 3 4 5 6 18 19 20 21 22 23
chain bonds :
 4-17 5-7 7-8 7-9 9-10 9-11 11-12 12-13 21-34 22-24 24-25 24-29 25-26 25-30 26-27
 27-28
ring bonds :
 1-2 1-6 2-3 3-4 4-5 5-6 18-19 18-23 19-20 20-21 21-22 22-23
exact/norm bonds :
 7-8 7-9 9-10 9-11 11-12 12-13 24-25 24-29 25-26 25-30 26-27 27-28
exact bonds :
 4-17 5-7 21-34 22-24
normalized bonds :
 1-2 1-6 2-3 3-4 4-5 5-6 18-19 18-23 19-20 20-21 21-22 22-23

```

G1:H,Ak

```

Match level :
 1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS
 12:CLASS 13:CLASS 17:CLASS 18:Atom 19:Atom 20:Atom 21:Atom 22:Atom 23:Atom 24:CLASS
 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS 34:CLASS

```

Generic attributes :

```

13:
Saturation : Saturated
Number of Carbon Atoms : less than 7
28:
Saturation : Saturated
Number of Carbon Atoms : less than 7

```

fragments assigned product role:
 containing 18

fragments assigned reactant/reagent role:
 containing 1

